```
3361 CCAGCT&ATTTTACTACAATCAGTAACGATAACGAGTATATGGATTA&CGCTGAAGCAGA 3420
Db
      3421 AAGTGGTGTGCATAATTCCTGAACATTTACAATCATACATTTCATCTTTATGGCGCCCAAA 3480
Qy
          Db
      3421 AAGTGGTGTGCATAATTCCTGAACATTTACAATCATACATTZCATCTTTATGGCGCCCAAA 3480
      3481 TAGTCATACTGTAAGAGGGGCTTTGCTGGATCTGCTTAAGGCTTCTTGTAAGTTGTG 3540
Qy
          3481 TAGTCATACTGTAAGAGGAGGCTTTGCTGGATCTGCTGTAAGGCTTCTTGTAAGTTGTG 3540
Db
      3541 GATGCCCCATTTTCTGGATGGGAACCTGCCAGACAGTGAACAAGGGCTTTGCAAGGTGCA 3600
Qу
          3541 GATGCCCCATTTTCTGGATGGGAACCTCCAGACAGTGAACAAGGGCTTTGCAAGGTGCA 3600
Db
      3601 GCATCCGGTTTTTGTTTTGCCAGTCCAAGAACCGTCCTCCTGTTACTTTGTAGTTGTACT 3660
Qy
          3601 GCATCCGGTTTTTGTTTTGCCAGTCCAAGAAACGCCTCCTGTTACTTTGTAGTTGTACT 3660
Db
      3661 CATACTAGTGCGCTTGTTTGTACAAGGAGAAATGTGTAACCTTGTTGAAAAAAATGTCTC 3720
Qy
      Db
Qy
      Db
      3781 CGAGAAATGGT#TTGTCGGTGTTAAGGTTGAAACGACTAGCTCTCGTTATCAAIGTGTTG 3840
Qy
          Db
      3781 CGAGAAAT GTTTTGTCGGTGTTAAGGTTGAAACGACTAGCTCTCGTTATCAATGTTTTG 3840
      3841 TAAACTTCTAGATTGATGTTTACCTTACTCTTGAAGTCAACACCGGAGAATTTAC 3~896
Qу
             Db
      3841 TAAACTTCTAGATTGATGTTTACCTTACTCTTGAAGTCAACACCGGAGAATTTAC 3896~
RESULT 2
US-10-447-135-1
; Sequence 1, Application US/10447135
Publication No. US20030199684A1
; GENERAL INFORMATION:
  APPLICANT: Hirochika, Hirohiko
  APPLICANT: Yamazaki, Muneo
  APPLICANT: Miyao, Akio
  TITLE OF INVENTION: A novel gene involved in brassinosteroid responses
  FILE REFERENCE: MAFF-1 DIV
  CURRENT APPLICATION NUMBER: US/10/447,135
  CURRENT FILING DATE: 2003-05-27
  PRIOR APPLICATION NUMBER: US 09/721,114
  PRIOR FILING DATE: 2000-11-22
  PRIOR APPLICATION NUMBER: JAPAN 2000-149106
  PRIOR FILING DATE: 2000-05-19
  NUMBER OF SEQ ID NOS: 3
  SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO 1
  LENGTH: 4310
  TYPE: DNA
  ORGANISM: Oryza sativa
  FEATURE:
  NAME/KEY: CDS
  LOCATION: (655)..(3828)
US-10-447-135-1
```

	cal :	Similarity 4; Conserva		Pred	d. :	No. 0	;	DB 6		Length Indels	4310;	Gaps	1;
Qy	1	CGCGGCTGTCG											60 .
Db	415	CGCGGCTGTCG											474
Qy	61	GATTGATTGTC											120
Db	475	GATTGATTGTC											534
Qy	121	TGTGCTTGGAT											180
Db	535	TGTGCTTGGAT		TGCT		TCGCG							594
Qy	181	TGAGTTGTGCC		GCTĞ	GCT	GTGTG							240
Db	595	TGAGTTGTGCC											654
Qy	241	ATGGAGATTGT											300
Db	655	ATGGAGATTGT											714
Qy	301	GCTCGTGGTGG.											360
Db	715	GCTCGTGGTGG											774
Qy	361	GATGCAGCCGC											420
Db	775	GATGCAGCCGC											834
Qy	421	ATAAGAGGGTA											480
Db	835	ATAAGAGGGTA											894
Qy	481	ATTTTCCATGA											540
Db	895	ATTTTCCATGA											954
Qy	541	GCAAAGTTTCG.											600
Db	955	GCAAAGTTTCG.	ACGATGGG	ATTG	CTC	GAAGT	GCT	TGGAT	'AA(GTTGAAA	ACTTCA	GATAAT	1014
Qy	601	GGAACAGCACC											660
Db	1015	GGAACAGCACC.											1074
Qy	661	ACATTTGTTCG											720
Db	1075	ACATTTGTTCG											1134
Qy	721	ACACAATCATC											780
Db	1135	ACACAATCATC	TCAAGGGA	AGAAT	rgc'	TGATA	GAT	CAACT	II CT:	TCCAAAG.	AGTGTG	CAAGAA	1194
Qy	781	GGCAATGACTC											840
Db	1195	GGCAATGACTC											1254

Qy	841	GATTCACCAATGAAAGATTTGCAAGGGCCAGCCCAAAATTATGATGTGGCAGCAAATGTC 90	0
Db	1255	GATTCACCAATGAAAGATTTGCAAGGGCCCAGCCCAAAATTATGATGTGGCAGCAAATGTC 13	14
Qy	901	TCTGAGGACAACACTTCTGTTGATGTTGGGGCTTTACCTGAAGTTCCCCAGATTACATGG 96	0
Db	1315	TCTGAGGACAACACTTCTGTTGATGTTGGGGCTTTACCTGAAGTTCCCCAGATTACATGG 13	74
Qy	961	CACATAGAAGTAAATGGTGCAGATCAACCTCCATCCACTCCAAAACTTTCTGAAGTGGTC 10	20
Db	1375	CACATAGAAGTAAATGGTGCAGATCAACCTCCATCCACTCCAAAACTTTCTGAAGTGGTC 14	34
Qy	1021	CTCAAAAGAAATGAAGATGAAAATGGAAAAACTGAAGAGACTCTTGTTGCTGAGCAGTGC 10	80
Db	1435	CTCAAAAGAAATGAAGATGAAAAATGGAAAAACTGAAGAGACTCTTGTTGCTGAGCAGTGC 14	94
Qу		AATTTGACCAAAGATCCTAACCCAATGTCTGGAAAGGAACGTGATCAGGTTGCTGAGCAG 11	
Db		AATTTGACCAAAGATCCTAACCCAATGTCTGGAAAGGAACGTGATCAGGTTGCTGAGCAG 15	
Qy		TGCAATTTGACCAAAGATCCGAAACCAGTGTCTGGGCAGAAATGTGAGCAGATCTGCAAT 12	
Db		TGCAATTTGACCAAAGATCCGAAACCAGTGTCTGGGCAGAAATGTGAGCAGATCTGCAAT 16	
Qy		GAGCCATGTGAAGAAGTTGTTCTCAAAAGAAGCTCCAAATCTAAGAGGAAGACGGATAAG 12	
Db		GAGCCATGTGAAGAAGTTGTTCTCAAAAGAAGCTCCAAATCTAAGAGGAAGACGGATAAG 16	
QУ		AAGTTGATGAAGAAGCAGCACAGCAAGAAACGCACTGCCCAGGCTGATGTTTCAGAT 13	
Db		AAGTTGATGAAGAAGCAGCACAGCAAGAAACGCACTGCCCAGGCTGATGTTTCAGAT 17	
Qy		GCAAAGCTTTGTCGGAGAAAGCCAAAAAAGGTGCGGCTTCTATCAGAAATTATAAATGCT 13	
Db		GCAAAGCTTTGTCGGAGAAAGCCAAAAAAGGTGCGGCTTCTATCAGAAATTATAAATGCT 17 AACCAGGTTGAGGATTCTAGAAGTGACGAAGTTCATCGTGAAAATGCCGCTGATCCCTGT 14	
Qy Db		AACCAGGTTGAGGATTCTAGAAGTGACGAAGTTCATCGTGAAAATGCCGCTGATCCCTGT 14	
Qy		GAGGATGATAGAAGTACCATCCCGGTCCCGATGGAAGTAAGCATGGATATTCCTGTTAGC 15	
Db		GAGGATGATAGAAGTACCATCCCGGTCCCGATGGAAGTAAGCATGGATATTCCTGTTAGC 19	
Qy		AACCATACAGTGGGAGAAGATGGGTTAAAATCAAGTAAGAACAAGACAAAACGCAAATAC 15	
Db			
Qy	1561	TCTGATGTTGTAGATGATCGATCACTTATGAACTGGCTGAATGGAAAAAAGAAAAGA 16	20
Db	1975		34
Qy	1621	ACTGGAAGTGTGCATCACACAGTTGCTCATCCAGCTGGGAATTTGAGCAACAAAAAAGTG 16	80
Db	2035		94
Qy	1681	ACACCCACTGCGAGTACTCAGCATGATGATGAGAATGATACTGAAAATGGTCTTGACACA 17	40
Db	2095		54
Qy	1741	AATATGCATAAGACAGATGTCTGTCAGCATGTATCAGAAATCTCCACACAGAGGTGCTCA 18	00

Db		
Qy	B01 TCAAAGGGGAAAACAGCGGGTTTGAGTAAGGGGAAAACACATTCAGCTGCTAGTAC	
Db		
Qy	361 TATGGTGGTGAAAGCACCAGAAATGGTCAGAACATACATGTACTCAGCGCAGAAGA	
Db		
Qy	921 TGCCAGATGGAAACCGAAAACTCTGTTCTGAGTCACTCGGCAAAGGTTTCTCCAGC	
Db		
Qy	981 CATGATATCCAAATTATGTCTGACCTTCATGAGCAGAGTCTACCCAAGAAGAAAAA	
Db		
Qy	041 CAAAAACTTGAAGTGACTCGTGAAAAACAGACCATGATAGATGACATCCCCATGGA	
Db		
Qy	01 GTTGAACTGCTAGCTAAAAACCAGCATGAGAGGCAGCTTATGACTGAGACTGATTG	
Db	515 GTTGAACTGCTAGCTAAAAACCAGCATGAGAGGCAGCTTATGACTGAGACTGATTG	
Qу	161 GACATCAACCGTATTCAATCCAAGACAACTGCTGATGATGATTGTGTAATAGTAGC	
Db	575 GACATCAACCGTATTCAATCCAAGACAACTGCTGATGATGATTGTGTAATAGTAGC	
Qу	221 AAGGATGGTTCAGATTATGCATCAAGTGTGTTTGACACTAATTCCCAACAGAAGTC	
Db	335 AAGGATGGTTCAGATTATGCATCAAGTGTGTTTGACACTAATTCCCAACAGAAGTC	
Qy	GCATCCCAAAGTACACAGAAGGAGTTACAGGGTCATTTGGCATTGACCACAAGA	
Db	595 GCATCCCAAAGTACACAGAAGGAGTTACAGGGTCATTTGGCATTGACCACACAAGA	
Qy	341 CCACATCCTCAGAACTTTCAGTCTACTCAGGAACAGCAGACACATTTGCGGATGGA	AGAA 2400
Db	755 CCACATCCTCAGAACTTTCAGTCTACTCAGGAACAGCAGACACATTTGCGGATGGA	AGAA 2814
Qy	ATGGTCACTATTGCTGCAAGCTCACCACTATTTTCACATCATGATGATCAGTATAT	
Db	315 ATGGTCACTATTGCTGCAAGCTCACCACTATTTTCACATCATGATGATCAGTATAT	
Qy	GAAGCACCAACTGAACATTGGGGCCGTAAGGACGCAAAGAAGCTAACGTGGGAGCA	
Db	375 GAAGCACCAACTGAACATTGGGGCCGTAAGGACGCAAAGAAGCTAACGTGGGAGCA	\ATTT 2934
Qy	521 AAGGCCACTACAAGAAATTCTCCAGCAGCAACATGTGGTGCTCAATTTAGACCTGG	
Db	335 AAGGCCACTACAAGAAATTCTCCAGCAGCAACATGTGGTGCTCAATTTAGACCTGG	
Qy	581 CAAGCAGTTGACTTGACTTCTACTCATGTCATGGGATCTTCCAGCAATTATGCATC	
Db	995 CAAGCAGTTGACTTGACTCATGTCATGGGATCTTCCAGCAATTATGCATC	TCGC 3054
Qу	541 CAACCAGTAATTGCGCCACTGGACCGCTATGCTGAAAGAGCGGTTAACCAGGTCCA	

Db	3055	${\tt CAACCAGTAATTGCGCCACTGGACCGCTATGCTGAAAGAGCGGTTAACCAGGTCCATGCA}$	3114
Qу	2701	AGAAATTTTCCAAGCACAATAGCAACCATGGAAGCGAGTAAGTTATGTGATCGGAGAAAT	2760
Db	3115	AGAAATTTTCCAAGCACAATAGCAACCATGGAAGCGAGTAAGTTATGTGATCGGAGAAAT	3174
Qу	2761	GCTGGACAAGTAGTCTTGTATCCTAAAGAATCCATGCCTGCGACGCATCTTCTGAGAATG	2820
Db	3175	GCTGGACAAGTAGTCTTGTATCCTAAAGAATCCATGCCTGCGACGCATCTTCTGAGAATG	3234
Qy	2821	ATGGATCCATCAACATTAGCAAGCTTCCCCAACTATGGAACTTCTAGCAGGAACCAGATG	2880
Db	3235	ATGGATCCATCAACATTAGCAAGCTTCCCCAACTATGGAACTTCTAGCAGGAACCAGATG	3294
Qy	2881	GAGTCTCAACTTCATAATTCTCAGTATGCACATAATCAGTACAAAGGATCAACCAGCACA	2940
Db	3295	GAGTCTCAACTTCATAATTCTCAGTATGCACATAATCAGTACAAAGGATCAACCAGCACA	3354
Qу	2941	TCATATGGCAGTAACCTGAATGGAAAGATTCCATTGACATTCGAAGACTTATCACGGCAT	3000
Ďb	3355	TCATATGGCAGTAACCTGAATGGAAAGATTCCATTGACATTCGAAGACTTATCACGGCAT	3414
Qу	3001	CAGCTGCATGATCTGCACAGACCTTTACGCCCACATCCTAGAGTTGGTGTGCTTGGCTCC	3060
Db	3415	CAGCTGCATGATCTGCACAGACCTTTACGCCCACATCCTAGAGTTGGTGTGTTGGCTCC	3474
Qy	3061	TTGCTGCAGAAGGAAATTGCAAACTGGTCGGAGAACTGTGGCACACAATCTGGTTATAAG	3120
Db	3475		3534
Qy	3121	TTAGGAGTGTCAACAGGAATAACATCGCATCAGATGAACAGAAAGGAACATTTTGAAGCC	3180
Db		TTAGGAGTGTCAACAGGAATAACATCGCATCAGATGAACAGAAAGGAACATTTTGAAGCC	3594
Qy	3181	CTGAATTCTGGAATGTTTTCAGCAAAATGGAATGCATTGCAGTTGGGTTCTGTTAGCTCC	3240
Db		CTGAATTCTGGAATGTTTTCAGCAAAATGGAATGCATTGCAGTTGGGTTCTGTTAGCTCC	
Qу	3241	AGTGCAGATTTTTTATCAGCGAGGAACAGCATAGCTCAATCTTGGACCAGAGGCAAGGGT	3300
Db	3655	AGTGCAGATTTTTTATCAGCGAGGAACAGCATAGCTCAATCTTGGACCAGAGGCAAGGGT	3714
Qy		AAAATGGTTCATCCCTTGGATCGGTTTGTGAGACAGGATATCTGTATAACTAAC	
Db		AAAATGGTTCATCCCTTGGATCGGTTTGTGAGACAGGATATCTGTATAACTAAC	
Qу		CCAGCTGATTTTACTACAATCAGTAACGATAACGAGTATATGGATTACCGCTGAAGCAGA	
Db		CCAGCTGATTTTACTACAATCAGTAACGATAACGAGTATATGGATTACCGCTGAAGCAGA	
Qу		AAGTGGTGTGCATAATTCCTGAACATTTACAATCATACATTTCATCTTTATGGCGCCAAA	
Db		AAGTGGTGTGCATAATTCCTGAACATTTACAATCATACATTTCATCTTTATGGCGCCAAA	
Qy		TAGTCATACTGTAAGAGGAGGGCTTTGCTGGATCTGCTGTAAGGCTTCTTGTAAGTTGTG	
Db		TAGTCATACTGTAAGAGGAGGGCTTTGCTGGATCTGCTGTAAGGCTTCTTGTAAGTTGTG	
Qy		GATGCCCCATTTTCTGGATGGGAACCTGCCAGACAGTGAACAAGGGCTTTGCAAGGTGCA	
Db	3955	GATGCCCCATTTTCTGGATGGGAACCTGCCAGACAGTGAACAAGGGCTTTGCAAGGTGCA	4014

```
3601 GCATCCGGTTTTTGTTTTGCCAGTCCAAGAAACGTCCTCCTGTTACTTTGTAGTTGTACT 3660
Qy
          4015 GCATCCGGTTTTTGTTTTGCCAGTCCAAGAAACGTCCTCCTGTTACTTTGTAGTTGTACT 4074
Db
Ov
      3661 CATACTAGTGCGCTTGTTTGTACAAGGAGAAATGTGTAACCTTGTTGAAAAAAATGTCTC 3720
          4075 CATACTAGTGCGCTTGTTTGTACAAGGAGAAATGTGTAACCTTGTTG-AAAAAATGTCTC 4133
Db
      Qy
          Db
      3781 CGAGAAATGGTTTTGTCGGTGTTAAGGTTGAAACGACTAGCTCTCGTTATCAATGTGTTG 3840
Qy
          Db
      4194 CGAGAAATGGTTTTGTCGGTGTTAAGGTTGAAACGACTAGCTCTCGTTATCAATGTGTTG 4253
      3841 TAAACTTCTAGATTGATGTGTTACCTTACTCTTGAAGTCAACACCGGAGAATTTAC 3896
Qу
          Db
      4254 TAAACTTCTAGATTGATGTTTACCTTACTCTTGAAGTCAACACCGGAGAATTTAC 4309
RESULT 3
US-10-437-963-40889
; Sequence 40889, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
 APPLICANT: La Rosa, Thomas J.
 APPLICANT: Kovalic, David K.
 APPLICANT: Zhou, Yihua
 APPLICANT: Cao, Yongwei
 APPLICANT: Wu, Wei
 APPLICANT: Boukharov, Andrey A.
 APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
 TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
 TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
 FILE REFERENCE: 38-21(53221)B
 CURRENT APPLICATION NUMBER: US/10/437,963
 CURRENT FILING DATE: 2003-05-14
 NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 40889
  LENGTH: 4640
  TYPE: DNA
  ORGANISM: Oryza sativa
 FEATURE:
  OTHER INFORMATION: Clone ID: PAT_MRT4530 44288C.1
US-10-437-963-40889
 Query Match
                  94.8%; Score 3693.4; DB 7; Length 4640;
 Best Local Similarity 99.9%; Pred. No. 0;
 Matches 3694; Conservative 0; Mismatches 1; Indels
       202 TGGCTGTGTTGATTCTCTCCTCGTCGTGGTGATCGATATGGAGATTGTTGCAGTAGAT 261
         Db
       946 TGCCTGTGTTGTTGATTCTCTCCTCGTCGTGGTGATCGATATGGAGATTGTTGCAGTAGAT 1005
      Db
```

WEST Search History

Hide Items Restore Clear Cancel

DATE: Monday, April 10, 2006

Hide?	Hit Count		
	DB=PC	SPB, USPT; PLUR=YES	; OP=ADJ
	L3	sung-zinmay-renee.in.	1
	L2	chen-lingjing.in.	2
	L1	moon-yong-hwan.in.	1

END OF SEARCH HISTORY

